# **Classification and Representation**

#### Classification

If we use linear regression, also y = 0 or 1.

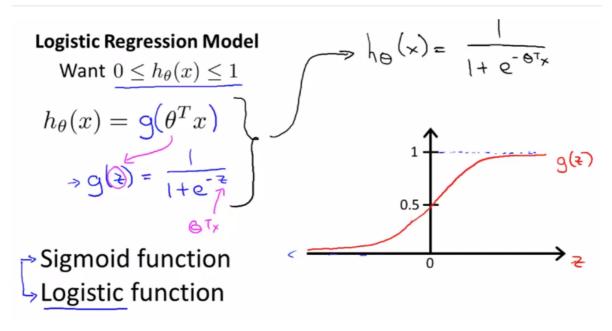
h(x) can be >1 or <0

So we consider using logistic regression: 0 < h(x) < 1

To attempt classification, one method is to use linear regression and map all predictions greater than 0.5 as a 1 and all less than 0.5 as a 0. However, this method doesn't work well because classification is not actually a linear function

The classification problem is just like the regression problem, except that the values we now want to predict take on only a small number of discrete values. For now, we will focus on the **binary classification problem** in which y can take on only two values, 0 and 1. (Most of what we say here will also generalize to the multiple-class case.) For instance, if we are trying to build a spam classifier for email, then  $x^{(i)}$  may be some features of a piece of email, and y may be 1 if it is a piece of spam mail, and 0 otherwise. Hence, ye $\{0,1\}$ . 0 is also called the negative class, and 1 the positive class, and they are sometimes also denoted by the symbols "-" and "+." Given  $x^{(i)}$ , the corresponding  $y^{(i)}$  is also called the label for the training example.

## **Hypothesis Representation**



### **Interpretation of Hypothesis Output**



 $h_{\theta}(x)$  = estimated probability that y = 1 on input x  $\leftarrow$ 

Example: If 
$$\underline{x} = \begin{bmatrix} x_0 \\ x_1 \end{bmatrix} = \begin{bmatrix} 1 \\ \text{tumorSize} \end{bmatrix}$$

$$\underline{h_{\theta}(x)} = \underline{0.7} \qquad \qquad \underline{y} = 0$$

Tell patient that 70% chance of tumor being malignant

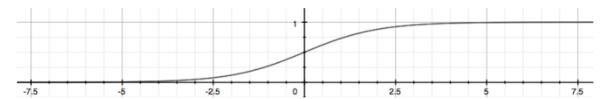
$$\frac{h_{\Theta}(x) = P(y=1|x;\theta)}{y=0} \qquad \text{"probability that } y=1, \text{ given } x, \\ \text{parameterized by } \theta"$$
 
$$\Rightarrow P(y=0|x;\theta) + P(y=1|x;\theta) = 1 \\ P(y=0|x;\theta) = 1 - P(y=1|x;\theta)$$

We could approach the classification problem ignoring the fact that y is discrete-valued, and use our old linear regression algorithm to try to predict y given x. However, it is easy to construct examples where this method performs very poorly. Intuitively, it also doesn't make sense for  $h_{\theta}(x)$  to take values larger than 1 or smaller than 0 when we know that y  $\in$  {0, 1}. To fix this, let's change the form for our hypotheses  $h_{\theta}(x)$  to satisfy  $0 \le h_{\theta}(x) \le 1$ . This is accomplished by plugging  $\theta^T x$  into the Logistic Function.

Our new form uses the "Sigmoid Function," also called the "Logistic Function":

$$egin{aligned} h_{ heta}(x) &= g( heta^T x) \ z &= heta^T x \ g(z) &= rac{1}{1 + e^{-z}} \end{aligned}$$

The following image shows us what the sigmoid function looks like:

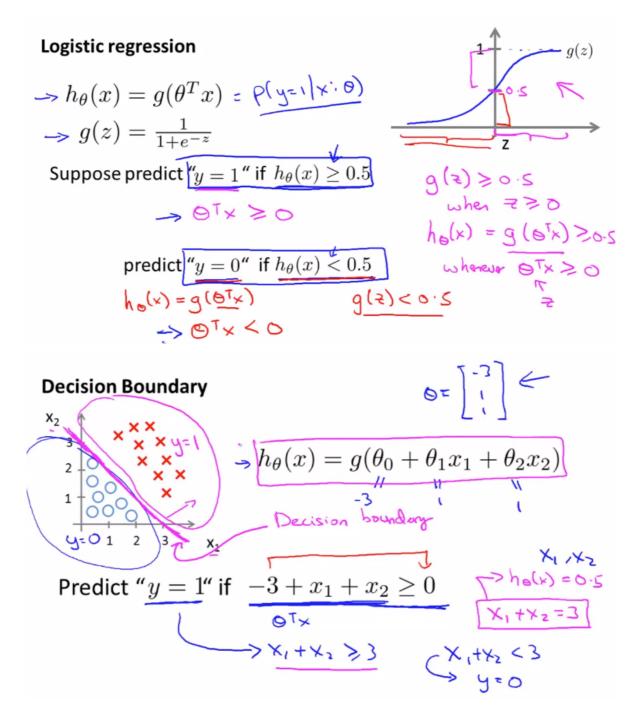


The function g(z), shown here, maps any real number to the (0, 1) interval, making it useful for transforming an arbitrary-valued function into a function better suited for classification.

 $h_{\theta}(x)$  will give us the **probability** that our output is 1. For example,  $h_{\theta}(x) = 0.7$  gives us a probability of 70% that our output is 1. Our probability that our prediction is 0 is just the complement of our probability that it is 1 (e.g. if probability that it is 1 is 70%, then the probability that it is 0 is 30%).

$$egin{aligned} h_{ heta}(x) &= P(y=1|x; heta) = 1 - P(y=0|x; heta) \ P(y=0|x; heta) + P(y=1|x; heta) = 1 \end{aligned}$$

### **Decision Boundaries**



The decision boundary is a property of the hypothesis. Even if we take away the data set, this decision boundary and the region where we predict y=1 versus y=0 that's a property of the hypothesis and of the parameters of the hypothesis and not a property of the data set.

In order to get our discrete 0 or 1 classification, we can translate the output of the hypothesis function as follows:

$$egin{aligned} h_{ heta}(x) &\geq 0.5 
ightarrow y = 1 \ h_{ heta}(x) &< 0.5 
ightarrow y = 0 \end{aligned}$$

The way our logistic function g behaves is that when its input is greater than or equal to zero, its output is greater than or equal to 0.5:

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g(z) \geq 0.5 \ when \ z \geq 0
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Remember.

$$egin{aligned} z=0,e^0=1&\Rightarrow g(z)=1/2\ z&\to\infty,e^{-\infty}&\to0\Rightarrow g(z)=1\ z&\to-\infty,e^\infty&\to\infty\Rightarrow g(z)=0 \end{aligned}$$

So if our input to g is  $\theta^T X$ , then that means:

$$h_{ heta}(x) = g( heta^T x) \geq 0.5 \ when \ heta^T x \geq 0$$

From these statements we can now say:

$$egin{aligned} heta^T x &\geq 0 \Rightarrow y = 1 \ heta^T x &< 0 \Rightarrow y = 0 \end{aligned}$$

The **decision boundary** is the line that separates the area where y = 0 and where y = 1. It is created by our hypothesis function.

#### Example:

$$egin{aligned} heta &= egin{bmatrix} 5 \ -1 \ 0 \end{bmatrix} \ y &= 1 \ if \ 5 + (-1)x_1 + 0x_2 \geq 0 \ 5 - x_1 \geq 0 \ -x_1 \geq -5 \ x_1 \leq 5 \end{aligned}$$

In this case, our decision boundary is a straight vertical line placed on the graph where  $x_1=5$ , and everything to the left of that denotes y = 1, while everything to the right denotes y = 0.

Again, the input to the sigmoid function g(z) (e.g.  $\theta^T X$ ) doesn't need to be linear, and could be a function that describes a circle (e.g.  $z=\theta_0+\theta_1x_1^2+\theta_2x_2^2$ ) or any shape to fit our data.